Seguences of the (+) strand

complete forms:

C12 C11 B12 E3 E12 A3 GGAGCUCAGACUUCACUCGUGUG---AGGCAAU-----AAAACA--UAAGUCCGAAGGUAACCAAUCCUAC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGCAAU-----AAAACG--UAAGUCCGAAGGUAACCAAUCCUAC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGCAAU-----AAAACUUGUAAGUCCGAAGGUAACCAAUCCUAC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGCAAUA-----AAAACU--UAAGUCCGAAGGUAACCAAUCCUAC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGGGGUGAGGCA------AAAAC--GUAAGACCGAAGGUAACCAUUCCUACCCACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGUAGUAAA----AAAAC--GUAAAUCCGAAGGUAACCAAUCCUAC-ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGUAGUAAAAA----AAAAC--GUAAAUCCGAAGGUAACCAAUCCUAC-ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGUAGUAAAAAA----AAAAC--GUAAAUCCGAAGGUAACCAGUCCUAC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGUAGUAAAAAAA----AAAAC--GUAAAUCCGAAGGUAACCAAUCCUAC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGUAGUAAAAAAA---AAAAC--GUAAAUCCGAAGGUAACCAAUCCUAC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGCAAU------AAAACU-UAAGUCCGAAGGUAACCAAUCCUGC--ACGUACCACUGUCGGUUCCAC GGAGCUCAGACUUCACU**CGUGUG---AGGCAAU--------AAAACU--UAAGUCCGAAGGUAACCAA**UCC**UAC--ACG**UACCACUGUCGGUUCCAC GGAGCUCAGACUUCACUCGUGUG---AGGCAGU-----AAAACU--UAAGUCCGAAGGUAACCAAUCCUAC--ACGUACCACUGUCGGUUCCAC

GGAGCUCAGACUUCACUCG DE.40F-Primer (Seq. ID No. 35) Primer moieties underlined and in bold

CGUACCACUGUCGGUUCCAC
DE.40R-Primer (rev. und
compl.) (Seq. ID No. 36)

Fig. 13-1

core rorms:		sed. In.	
CGUGUGAGGCAAUAAAACUUAAGUCCGAAGGUAACCAAUCCUACACG	B11	<b>&amp;</b>	
CGUGUGAGGCAGUAAACUUAAGUCCGAAGGUAACCAAUCCUACACG	<b>G</b> 2	10	
CGUGUGAGCCAAUAAACUUAAGUCCGAAGGUAACCAAUCCUGCACG	E12	12	
CGUGUGAGGCAAUAAACAUAAGUCCGAAGGUAACCAAUCCUACACG	B7	14	
CGUGUGAGGCAAUAAAACGUAAGUCCGAAGGUAACCAAUGCUACAGG	A8	16	:
CGUGUGAGGCAAUAAAACUUGUAAGUCCGAAGGUAACCAAUCCUACACG	B12	8	
CGUGUGAGGCAAUAAAACUUAAGUCCGAAGGUAACCAAUCCUACACG	E3	20	
CGUGCGGUGAGGCAAAAACGUAAGACCGAAGGUAACCAUUCCUACCCACG	C12	22	
CGUGUGAGGUAGUAAAAAAACGUAAAUCCGAAGGUAACCAAUCCUACACG	C11	24	
CGUGUGAGGUAGUAAAAAAAAAACGUAAAUCCGAAGGUAACCAAUCCUACACG	A3	26	
CGUGUGAGGUAGUAAAAAAAAAACGUAAAUCCGAAGGUAACCAGUCCUACACG	FS	28	
CGUGUGAGGUAGUAAAAAAAAAAAAGGUAAAGGUAACCAAUCCUACACG	A12	30	
CGUGUGAGGUAGUAAAAAAAAAAAAACGUAAAUCCGAAGGUAACCAAUCCUACACG	F12	32	
CGUGUGAGGAGUAAAAAAAAAAAAAAAAAAAAAAAAAAA	<b>G</b> 2	34	

ig. 13 -2

Sequences derived from the RNA selection (round 13)

Sed	Seq. RNA round 13	
ID.	group1	
38	SOT-RO4-DR13-ES	GGAGCTCAGACTTCACTCGTGCGGTGAGGCAGACG-TAAGACCGAAGGTAACCATTCCTACCCACGTACCACTTCGGTTCCAC
	group2 2.1 main clone	
တ္ထ	*SOT-R04~DR13-A2	GGAGCTCAGACTTCACTCGTGCGGTGAGGCTAACG-TAAGACCGAAGGTAACCATTCCTACCCACGTACCACTGTCGGTTCCAC
40	variations of 2.1 'SOT-R04-DR13-C4	GGAGCTCAGACTTCACTCGTGTGGTGAGGCTAACG-TAAGACCGAGGTAACCATTCCTACCCACGTACCACTGTCGGTTCCAC
	group3	
41	'SOT-R04-DR13-C1	GGAGCTCAGACTTCACTCGTGTGAGGTAATAAAAC-TAAATCCGAAGGTAACCAATCCTACACGTACCACTGTCGGTTCCAC
	group4	
42	'SOT-R04-DR13-G2	GGAGCTCAGACTTCACTCGTGCGTACCGTTATAAAGGGAGTCCTGCAGACTGATGCCAGGCCAGGTACCACTGTCGGTTCCAC

Fig. 22

Sequences derived from the RNA selection (round 14)

Seg	Seq. RNA round 14	
ID.	group1 1.1 (main clone)	
43	'SOT-R04-DR14-F7	GGAGCTCAGACTTCACTCGTGCGGTGAGGCAGACGTAAGACCGAAGGTAACCATTCCTACCAGGTACCACTGTCGGTTCCAC
	variations of clone 1.2	clone 1.1
44	'SOT-R04-DR14-C11	GGAGCTICAGACTTCACTCGTGCGGTGAGGCAAACGTAAGACCGAAGGTAACCATTCCTACCACGTACCACTGTCGGTTCCAC
45	'SOT-R04-DR14-A8	GGAGCTCAGACTTCACTCGTGCGGTGAGGCTAACGTAAGACCGAAGGTAACCATTCCTACCACGTACCACTGTCGGTTCCAC
46	*SOT-R04-DR14-C12	GGAGCTCAGACTTCACTCGTGCGGTGAGGCAAAAACGTAAGACCGAAGGTAACCATTCCTACCACGTACCACTGCGGTTCCAC

48 'SOT-R04-DR14-E11 GGAGCTCAGACTICACTCGTGTGAGGCAGTAAACTTAAGTCCGAAGGTAACCAATCCTAACGTACCACTGTGGGTTCCAC  1.3 49 'SOT-R04-DR14-H11 GGAGCTCAGACTTCACTCGTGTGAGGCAATTAAACTTG-AAGTCCGAAGGTAACCAATCCTAACGTACCACTGTGGGTTCCAC  1.4
variations of clone 3.1

Sequences derived from the 2 F-RNA selection (round 14)

7	circos acritoca troll	
Seq.	Seq. 2 -F-RNA rou	round 14
ID.	group1	
51	1.1 (main clone) 'SOT-F03-DR14-G6	GGAGCICAGACITICACICGIGGAATAGGAATGACTCAGACGTITCT—CATACGTCGCCGCAACGIACCACTGTCGGTICCAC
	mutations of clone 1.1	

	mutations of clone 1.1	
	7.7	
52	.sor-F03-DR14-F2	GGAGCTCAGACTICACTCGTGGAATAGGAATGACTCAGACCTTTCTCATAGGTCGCCGCAACGTACCACTGTCGGTTCCAC
53	'SOT-F03-DR14-F4.	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACCTTTCTATAGGTCGCCGCAACGTACCACTGTCGGTTCCAC
54	'SOT-F03-DR14-D5	GGAGCTCAGACTTCACTCGGATAGGATGACTCAGACGTTTCTC-CATACGTCGCCGCGACGTACCACTGTCGGTTCCAC
55	'SOT-F03-DR14-G3	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTTTCTC-CAAACGTCGCCGCAACGTACCACTGTCGGTTCCAC
26	'SOT-F03-DR14-B5	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTT-CTC-CATACGTCGCCGCAACGTACCACTGTCGGTTCCAC
57	'SOT-F03-DR14-C2	GGAGCTCAGACTICACTCGTGGAATAGGAATGACTCAGACCTTTCTC-CATAGGTCGCCAACGTACCACTGTCGGTTCCAC
58	'SOT-F03-DR14-F3	GGAGCTCAGACTICACTCGTGGAATAGGAATGACTCAGACGTTTCTT-CATACGTCGCCGCAACGTACCACTGTCGGTTCCAC
59	'SOT-F03-DR14-B6	GGAGCTCAGACTTCACTCGTGGAATAGGAATGATTCAGACGTTTCCATACGTCGCCGCAACGTACCACTGTCGGTTCCAC
09	'SOT-F03-DR14-H1	GGAGCTCAGACTICACTCGTGGAATAGGAATGACTCAGACCTTTCATAGGTCGCCGCAACGTACCACTGTCGGTTCCAC
19	'SOT-F03-DR14-F6 1.13	GGAGCTCAGACTTCACTCGTGCAATAGGAATGACTCAGACGTTTCATACGTCGCCGCAACGTACCACTGTGGGTTCCAC
62	'SOT-F03-DR14-B1	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTCCATACGTCGCCGCAACGTACCACTGTCGGTTCCAC
63	'SOT-F03-DR14-C1	GGAGCTCAGACTTCACTCGTGCAATAGGAATGACTCAGACGTTTTCTCCCATACGTCGCCGCAACGTACCACTGTGGGTTCCAC
64	*SOT-F03-DR14-H5	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTTTTTC-CATACGTCGCCGCAACGTACCACTGTCGGTTCCAC

GGAGCTCAGACTTCACTCGTGCGCTTTCTGTTAGCTGCCGACOGTCAGTGCGGCACGAGATACGTACGTACGTGTCGGTTTCCAC	clone 2.1	GGAGCTCAGACTTCACTGGTGCGCTTTCTGTTAGCTGCTGACCGTCAGTGCGGCACGAGATACGTACCACTGTCGGTTCCAC	GGAGCTCAGACTTCACTCGTGCGCTTTCTGTTAGCT-CCGACCGTCAGTGCGGCACGAGATACGTACGACTGTCGGTTCCAC	GGAGCTCAGACTTCACTCGTGCGCTTTCTGTTAGCT -CAGACCGTCAGTGCGGCACGAGATACGTACCACTGTCGGTTCCAC	GGAGCTCAGACTTCACTCGTGCGCTTT-TGTTAGCT-CAGACCGTCAGTGCGGCACGAGATACGTACCACTGTCGGTTCCAC	ggagctcagacttcactcgtgcgctttctagctcttaaccgaccgtgcggcacgag <u>acgtaccactgtcggttccac</u>	GGAGCTCAGACTTCACTCGTGTGCCGCCCCTTATTGTCAGGGAGCTTGAGCCGACGACGTACCACTGTCGGTTCCAC
Seq. <b>group2</b> ID. 2.1 (main clone) 65 'SOT-F03-DR14-G5	variations of clos	'SOT=F03-DR14-D3	'SOT-F03-DR14-H2	'SOT-F03-DR14-D1	'SOT-F03-DR14-A2	.SOT-F03-DR14-G2	group3 3.1 main clone 'SOT-F03-DR14-H6
Seq. ID. 65		99	19	89	69	70	11

Fig. 24-2:

Sedn	ences derived from	Sequences derived from the 2'F-RNA selection (round 15)
Seg.	Seq. 2 -F-RNA rou	ound 15
ID.	group1	
	1.1 (main clone)	
72	'SOT-F03-DR15-G10	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGAGGTTTCT-CATAGGT-CGCCGCAACGTACCACTGTCGGTTCCAC
73	*:2 *SOT-F03-DR15-G7	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTTTCT-CAAACGT-CGCCGCAACGTACCACTGTCGACTTCCAC
74	3 .sor-F03-DR15-F10	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTTTCCATACGT-CGCCGCAACGTACCACTGTCGGTTCCAC
75	1.4 'SOT-F03-DR15-D9	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTTTCATGT-CGCCGCAACGTACCACTGTCGGTTCCAC
76	SOT-F03-DR15-F12	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTTTCATACGT-CGCCGCAACGTACCACTGTCGGTTCCAC GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTTTCATACGT-CGCCGCAACGTACCACTGTCGGTTCAC
78	1.6 'SOT-F03-DR15-H7	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACGTCCATACGT-CGCCGCAACGTACCACTGTCGGTTCCAC
79	SOT-F03-DR15-A11	GGAGCTCAGACTTCACTOGTGGAATAGGAATGACTCAGACCTTTCCATAGGT-CGCCGCAACGTACCACTGTCGGTTCCAC
. 80	SOT-F03-DR15-A8	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACCTTTCT-CATAGGT-CGCCGCA <u>ACGTACCACTGTCGGTTCCAC</u>
81	1.10	<u>ggagctcagacttcagtcgtg</u> gaataggaatgactcagacgtttctccatacgt-cgccgca <u>acgtaccactgtcggttccac</u>
82	'SOT-F03-DR15-C9	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGGCCTTTCTTCATAGGT-CGCCGCAACGTACCACTGTCGGTTCCAC
83	'SOT-F03-DR15-C12	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACCCAGACGTTTTCATACGT-CGCCGCA <u>ACGTACCACTGTCGGTTCCAC</u>
84	'SOT-F03-DR15-F7	<u> ggagctcagacttcactcoto</u> gaataggaatgactcagaccttttt-cataggt-cgccgca <u>acgtaccactotoggttccac</u>
85	'SOT-F03-DR15-A12	GGAGCTCAGACTTCACTCGTGGAATAGGAATGACTCAGACCTTTTT-CATAGGT-CGCCGCAACGTACCACTGTCGGTTCCAC

86

group2
2.1
'SOT-F03-DR15-C7